



SEQUENCE LISTING

<110> Feussner et al.

<120> Linoleate- and Linolenate-Lipoxygenase-Mutants

<130> 213721

<141> US 09/937,908

<140> 2002-01-07

<150> PCT/EP00/02545

<151> 2000-03-22

<150> DE 19914464.8

<151> 1999-03-30

<160> 5

<170> PatentIn Ver. 3.2

<210> 1

<211> 878

<212> PRT

<213> Cucumis sativus

<400> 1

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Arg Val Ser Ser Leu Gly Gly Asn Lys Ile Lys Gly Lys Val Ile Leu
35 40 45

Met Arg Ser Asn Val Leu Asp Phe Thr Glu Phe His Ser Asn Leu Leu
50 55 60

Asp Asn Phe Thr Glu Leu Leu Gly Gly Gly Val Ser Phe Gln Leu Ile
65 70 75 80

Ser Ala Thr His Thr Ser Asn Asp Ser Arg Gly Lys Val Gly Asn Lys
85 90 95

Ala Tyr Leu Glu Arg Trp Leu Thr Ser Ile Pro Pro Leu Phe Ala Gly
100 105 110

Glu Ser Val Phe Gln Ile Asn Phe Gln Trp Asp Glu Asn Phe Gly Phe
115 120 125

Pro Gly Ala Phe Phe Ile Lys Asn Gly His Thr Ser Glu Phe Phe Leu
130 135 140

Lys Ser Leu Thr Leu Asp Asp Val Pro Gly Tyr Gly Arg Val His Phe
145 150 155 160

Asp	Cys	Asn	Ser	Trp	Val	Tyr	Pro	Ser	Gly	Arg	Tyr	Lys	Lys	Asp	Arg	165	170	175
Ile	Phe	Phe	Ala	Asn	His	Val	Tyr	Leu	Pro	Ser	Gln	Thr	Pro	Asn	Pro	180	185	190
Leu	Arg	Lys	Tyr	Arg	Glu	Glu	Glu	Leu	Trp	Asn	Leu	Arg	Gly	Asp	Gly	195	200	205
Thr	Gly	Glu	Arg	Lys	Glu	Trp	Asp	Arg	Ile	Tyr	Asp	Tyr	Asp	Val	Tyr	210	215	220
Asn	Asp	Ile	Ala	Asp	Pro	Asp	Val	Gly	Asp	His	Arg	Pro	Ile	Leu	Gly	225	230	235
Gly	Thr	Thr	Glu	Tyr	Pro	Tyr	Pro	Arg	Arg	Gly	Arg	Thr	Gly	Arg	Pro	245	250	255
Arg	Ser	Arg	Arg	Asp	His	Asn	Tyr	Glu	Ser	Arg	Leu	Ser	Pro	Ile	Met	260	265	270
Ser	Leu	Asp	Ile	Tyr	Val	Pro	Lys	Asp	Glu	Asn	Phe	Gly	His	Leu	Lys	275	280	285
Met	Ser	Asp	Phe	Leu	Gly	Tyr	Thr	Leu	Lys	Ala	Leu	Ser	Ile	Ser	Ile	290	295	300
Lys	Pro	Gly	Leu	Gln	Ser	Ile	Phe	Asp	Val	Thr	Pro	Asn	Glu	Phe	Asp	305	310	315
Asn	Phe	Lys	Glu	Val	Asp	Asn	Leu	Phe	Glu	Arg	Gly	Phe	Pro	Ile	Pro	325	330	335
Phe	Asn	Ala	Phe	Lys	Thr	Leu	Thr	Glu	Asp	Leu	Thr	Pro	Pro	Leu	Phe	340	345	350
Lys	Ala	Leu	Val	Arg	Asn	Asp	Gly	Glu	Lys	Phe	Leu	Lys	Phe	Pro	Thr	355	360	365
Pro	Glu	Val	Val	Lys	Asp	Asn	Lys	Ile	Gly	Trp	Ser	Thr	Asp	Glu	Glu	370	375	380
Phe	Ala	Arg	Glu	Met	Leu	Ala	Gly	Pro	Asn	Pro	Leu	Leu	Ile	Arg	Arg	385	390	395
Leu	Glu	Ala	Phe	Pro	Pro	Thr	Ser	Lys	Leu	Asp	Pro	Asn	Val	Tyr	Gly	405	410	415
Asn	Gln	Asn	Ser	Thr	Ile	Thr	Glu	Glu	His	Ile	Lys	His	Gly	Leu	Asp	420	425	430
Gly	Leu	Thr	Val	Asp	Glu	Ala	Met	Lys	Gln	Asn	Arg	Leu	Tyr	Ile	Val	435	440	445
Asp	Phe	His	Asp	Ala	Leu	Met	Pro	Tyr	Leu	Thr	Arg	Met	Asn	Ala	Thr	450	455	460

Ser Thr Lys Thr Tyr Ala Thr Arg Thr Leu Leu Leu Leu Lys Asp Asp
 465 470 475 480
 Gly Thr Leu Lys Pro Leu Val Ile Glu Leu Ala Leu Pro His Pro Gln
 485 490 495
 Gly Asp Gln Leu Gly Ala Ile Ser Lys Leu Tyr Phe Pro Ala Glu Asn
 500 505 510
 Gly Val Gln Lys Ser Ile Trp Gln Leu Ala Lys Ala Tyr Val Thr Val
 515 520 525
 Asn Asp Val Gly Tyr His Gln Leu Ile Ser His Trp Leu His Thr His
 530 535 540
 Ala Val Leu Glu Pro Phe Val Ile Ala Thr His Arg Gln Leu Ser Val
 545 550 555 560
 Leu His Pro Ile His Lys Leu Leu Val Pro His Tyr Lys Asp Thr Met
 565 570 575
 Phe Ile Asn Ala Ser Ala Arg Gln Val Leu Ile Asn Ala Asn Gly Leu
 580 585 590
 Ile Glu Thr Thr His Tyr Pro Ser Lys Tyr Ser Met Glu Leu Ser Ser
 595 600 605
 Ile Leu Tyr Lys Asp Trp Thr Phe Pro Asp Gln Ala Leu Pro Asn Asn
 610 615 620
 Leu Met Lys Arg Gly Leu Ala Val Glu Asp Ser Ser Ala Pro His Gly
 625 630 635 640
 Leu Arg Leu Leu Ile Asn Asp Tyr Pro Phe Ala Val Asp Gly Leu Asp
 645 650 655
 Ile Trp Ser Ala Ile Lys Thr Trp Val Gln Asp Tyr Cys Cys Leu Tyr
 660 665 670
 Tyr Lys Asp Asp Asn Ala Val Gln Asn Asp Phe Glu Leu Gln Ser Trp
 675 680 685
 Trp Asn Glu Leu Arg Glu Lys Gly His Ala Asp Lys Lys His Glu Pro
 690 695 700
 Trp Trp Pro Lys Met Gln Thr Leu Ser Glu Leu Ile Glu Ser Cys Thr
 705 710 715 720
 Thr Ile Ile Trp Ile Ala Ser Ala Leu His Ala Ala Val Asn Phe Gly
 725 730 735
 Gln Tyr Pro Tyr Gly Gly Tyr Ile Leu Asn Arg Pro Thr Thr Ser Arg
 740 745 750
 Arg Phe Met Pro Glu Val Gly Thr Ala Glu Tyr Lys Glu Leu Glu Ser
 755 760 765

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Asn Pro Glu Lys Ala Phe Leu Arg Thr Ile Cys Ser Glu Leu Gln Ala
770 775 780

Leu Val Ser Ile Ser Ile Ile Glu Ile Leu Ser Lys His Ala Ser Asp
785 790 795 800

Glu Val Tyr Leu Gly Gln Arg Ala Ser Ile Asp Trp Thr Ser Asp Lys
805 810 815

Ile Ala Leu Glu Ala Phe Glu Lys Phe Gly Lys Asn Leu Phe Glu Val
820 825 830

Glu Asn Arg Ile Met Glu Arg Asn Lys Glu Val Asn Leu Lys Asn Arg
835 840 845

Ser Gly Pro Val Asn Leu Pro Tyr Thr Leu Leu Val Pro Ser Ser Asn
850 855 860

Glu Gly Leu Thr Gly Arg Gly Ile Pro Asn Ser Ile Ser Ile
865 870 875

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<213> Artificial

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<211> 59
<212> DNA
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BEST AVAILABLE COPY

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<210> 5

<211> 57

<212> DNA

<213> Artificial

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<223> Oligonucleotide primer

<400> 5

gtatgcaacc aatgactaat aagttgatgg taaccgaaat cattaacagt tacataa 57